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## IN THE CLAIMS

- 29. (Amended) Computer readable medium having recorded thereon at least 100 of the nucleotide sequences depicted in SEQ ID NO: 16207 through SEQ ID NO: 27905 or complements thereof.
- 30. (Amended) Computer readable medium according to claim 29, having recorded thereon at least 1000 said nucleotide sequences.
- 47. (New) Computer readable medium of claim 29 wherein said nucleotide sequences are selected from the group consisting of SEQ ID NO: 16207 through SEQ ID NO: 26804 and complements thereof.
- 48. (New) Computer readable medium of claim 29 wherein said nucleotide sequences are selected from the group consisting of SEQ ID NO: 16207 through SEQ ID NO: 25999 and complements thereof
- 49. (New) Computer readable medium of claim 29 wherein said nucleotide sequences are selected from the group consisting of SEQ ID NO: 16207 through SEQ ID NO: 24034 and complements thereof
- 50. (New) Computer readable medium of claim 29 wherein said nucleotide sequences are selected from the group consisting of SEQ ID NO: 16207 through SEQ ID NO: 22709 and complements thereof.
- 51. (New) Computer readable medium of claim 29 wherein said nucleotide sequences are selected from the group consisting of SEQ ID NO: 16207 through SEQ ID NO: 17680 and complements thereof.

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- 52. (New) Computer readable medium of claim 29 wherein said nucleotide sequences are selected from the group consisting of SEQ ID NO: 16207 through SEQ ID NO: 17617 complements thereof.
- 53. (New) Computer readable medium of claim 29 wherein said nucleotide sequences are selected from the group consisting of SEQ ID NO: 16207 through SEQ ID NO: 17294 and complements thereof.
- 54. (New) Computer readable medium comprising the nucleotide sequences of the group consisting of SEQ ID NO: 1620 through SEQ ID NO: 26804.
- 55. (New) Computer readable medium of claim 29 which is adapted for homology searching of nucleotide sequence.
- 56. (New) Computer readable medium of claim 55 wherein said medium is adapted for homology searching using a BLAST algorithm.
- 57. (New) A computer based system comprising computer readable medium of claim 29, input means for receiving a target sequence, means for identifying fragments of sequence recorded in said computer readable medium which are homologous to a target sequence, and an output means for outputting identified homologous sequences.
- 58. (New) A method of identifying nucleotide sequence comprising comparing target sequence to a sequence stored in computer readable medium of claim 29.

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